```
============================================================
Sequence Listing could not be accepted due to errors.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: Keisha Douglas
Timestamp: [year=2010; month=11; day=3; hr=10; min=53; sec=1; ms=998; ]
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Reviewer Comments:
<210>  1

<211>  7

<212>  PRT

<213>  Gymnea sylvestre



<400>  1



Asn Gly Ser Phe Ser Gly Phe

1               5


 The above sequence id# 1 is invalid, please delete spacing betweeen the
amino acids and the numbering.   This error is seen globally throughout
the sequences.
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Application No: 10579655   Version No: 4.0

Input Set:

Output Set:

Started:	2010-10-25 19:40:21.780
Finished:	2010-10-25 19:40:36.460
Elapsed:	0 hr(s) 0 min(s) 14 sec(s) 680 ms
Total Warnings:	379
Total Errors:	2984
No. of SeqIDs Defined:	415
Actual SeqID Count:	415

Error code		Error Description
W	402	Undefined organism found in <213> in SEQ ID (1)
E	355	Empty lines found between the amino acid numbering and the
E	321	No. of Bases conflict, this line has no nucleotides SEQID (1)
W	402	Undefined organism found in <213> in SEQ ID (2)
E	355	Empty lines found between the amino acid numbering and the
E	321	No. of Bases conflict, this line has no nucleotides SEQID (2)
W	402	Undefined organism found in <213> in SEQ ID (3)
E	355	Empty lines found between the amino acid numbering and the
E	321	No. of Bases conflict, this line has no nucleotides SEQID (3)
W	402	Undefined organism found in <213> in SEQ ID (4)
E	355	Empty lines found between the amino acid numbering and the
E	321	No. of Bases conflict, this line has no nucleotides SEQID (4)
W	213	Artificial or Unknown found in <213> in SEQ ID (5)
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E	321	No. of Bases conflict, this line has no nucleotides SEQID (5)
E	355	Empty lines found between the amino acid numbering and the
E	321	No. of Bases conflict, this line has no nucleotides SEQID (5)
W	213	Artificial or Unknown found in <213> in SEQ ID (6)
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E	321	No. of Bases conflict, this line has no nucleotides SEQID (6)

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                    Input Set:


                    Output Set:


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                    Finished:   2010-10-25 19:40:36.460
                     Elapsed:   0 hr(s) 0 min(s) 14 sec(s) 680 ms
              Total Warnings:   379
                Total Errors:   2984
         No. of SeqIDs Defined: 415
             Actual SeqID Count: 415
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Error code	Error Description
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E 321	No. of Bases conflict, this line has no nucleotides SEQID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
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E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
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E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (8)
W 402	Undefined organism found in <213> in SEQ ID (9)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (9)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (9)
W 402	Undefined organism found in <213> in SEQ ID (10)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (10)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (10)

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                    Input Set:


                    Output Set:


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                    Finished:   2010-10-25 19:40:36.460
                     Elapsed:   0 hr(s) 0 min(s) 14 sec(s) 680 ms
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E   321         No. of Bases conflict, this line has no nucleotides  SEQID (11)

E   355         Empty lines found between the amino acid numbering and the

E   321         No. of Bases conflict, this line has no nucleotides  SEQID (11)

E   355         Empty lines found between the amino acid numbering and the

E   321         No. of Bases conflict, this line has no nucleotides  SEQID (11)

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E   355         Empty lines found between the amino acid numbering and the
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E   321         No. of Bases conflict, this line has no nucleotides  SEQID (12)
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W   213         Artificial or Unknown found in <213>  in SEQ ID (19)

W   213         Artificial or Unknown found in <213>  in SEQ ID (20)

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          Total Errors:   2984
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      Actual SeqID Count: 415


Error code      Error Description

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W   402         Undefined organism found in <213>  in SEQ ID (30)
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W   213         Artificial or Unknown found in <213>  in SEQ ID (77)

W   213         Artificial or Unknown found in <213>  in SEQ ID (78)

W   213         Artificial or Unknown found in <213>  in SEQ ID (79)

W   213         Artificial or Unknown found in <213>  in SEQ ID (80)

W   213         Artificial or Unknown found in <213>  in SEQ ID (253)

W   213         Artificial or Unknown found in <213>  in SEQ ID (254)

W   213         Artificial or Unknown found in <213>  in SEQ ID (255)

W   213         Artificial or Unknown found in <213>  in SEQ ID (256)

W   213         Artificial or Unknown found in <213>  in SEQ ID (257)

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<110> Sanofi Pasteur, Inc.

<120> METHODS FOR PURIFYING PERTUSSIS TOXIN AND PEPTIDES USEFUL

       THEREFOR

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<140> 10579655
<141> 2010-10-25

<150> 60/523,881

<151> 2003-11-20

<150> PCT/US2004/038700

<151> 2004-11-18

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Asn Met Arg Ile Glu Thr Pro Asn Asn Ile Arg Lys Asp Ala

          20   25   30

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            20   25   30

<210> 8
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<400> 8

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<211> 34

<212> PRT

<213> Gymnea sylvestre

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Ser Gly Cys Val Lys Lys Asp Glu Leu Cys Ala Arg Trp Asp Leu Val

1   5   10   15

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            20   25   30

Cys Gly

<210> 10

<211> 34

<212> PRT

<213> Gymnea sylvestre

<400> 10

Ser Gly Cys Val Lys Lys Asp Glu Leu Cys Glu Leu Ala Val Asp Glu

1   5   10   15

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Cys Gly

<210> 11

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<212> PRT

<213> Gymnea sylvestre

<400> 11

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Cys Cys Glu Pro Leu Glu Cys Lys Trp Phe Asn Glu Asn Tyr Gly Ile
            20   25   30

Cys Gly Ser
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<213> Gymnea sylvestre

<400> 12

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            20   25   30

Cys Gly

<210> 13

<211> 35

<212> PRT

<213> Gymnea sylvestre

<400> 13

Gln Gln Cys Val Lys Lys Asp Glu Leu Cys Ile Pro Tyr Tyr Leu Asp

1   5   10   15

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        20   25   30

Cys Ile Gly

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<210> 14

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nnsnnstgct gtgagcccct cgagtgcnns nnsnnsnnsn nsnnsnnsnn snnstgcggc   120

agcggcagtt ctgggtctag c   141

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<213> Gymnea sylvestre

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catagtggct caagctcagg atca   84

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<210> 19

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        20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Asp Ala Asn Ala Pro

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Lys Ala Ser Ala Ile

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His His His His His His

1   5

<210> 21

<211> 6

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<213> Artificial Sequence

<220>

<223> Synthetic sequence, no source organism

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1   5

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<223> Synthetic sequence, no source organism

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snnsnnsnns nnsnnsnnsn nsnnsnnsnn snnsnnsnns nnsnnagatc tagcatgatg   120

atgatga   127

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<211> 81

<212> DNA

<213> Gymnea sylvestre

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catcatcatc atgctagatc t   81

<210> 24

<211> 32

<212> DNA

<213> Gymnea sylvestre

<400> 24

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<210> 25

<211> 18

<212> DNA

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<400> 25

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<213> Gymnea sylvestre

<400> 26

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1   5   10   15

Val Lys Lys Asp Glu Leu Cys Ala Gly Ser Val Gly His Cys Cys Glu
          20   25   30

Pro Leu Glu Cys Leu Arg Arg Phe Leu Asn Leu Arg Trp Cys Gly Ser
          35   40   45

Gly Ser Ser Gly Ser Ser
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<210> 27

<211> 54

<212> PRT

<213> Gymnema sylvestre

<400> 27

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys
1   5   10   15

Val Lys Lys Asp Glu Leu Cys Ile Val Met Arg Ala Pro Cys Cys Glu
          20   25   30

Pro Leu Glu Cys Leu Arg Arg Tyr Met Leu Lys His Met Cys Gly Ser

        35   40   45

Gly Ser Ser Gly Ser Ser

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<210> 28

<211> 54

<212> PRT

<213> Gymnea sylvestre

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Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1   5   10   15

Val Lys Lys Asp Glu Leu Cys Lys Ala Phe Arg Tyr Ser Cys Cys Glu

            20   25   30

Pro Leu Glu Cys Leu Arg Lys Trp Leu Lys Ala Arg Phe Cys Gly Ser

        35   40   45

Gly Ser Ser Gly Ser Ser

    50

<210> 29

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<213> Gymnea sylvestre

<400> 29

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1   5   10   15

Val Lys Lys Asp Glu Leu Cys Leu Arg Ser Ser Ile Asp Cys Cys Glu

            20   25   30

Pro Leu Glu Cys Leu Tyr Lys Trp Met Gln Arg Arg Leu Cys Gly Ser

        35   40   45

Gly Ser Ser Gly Ser Ser

    50

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<210> Gymnea sylvestre

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1   5   10   15

Val Lys Lys Asp Glu Leu Cys Trp Pro Arg Arg His Lys Cys Cys Glu
            20   25   30

Pro Leu Glu Cys Leu Leu Glu Met Leu Glu Arg Lys Arg Cys Gly Ser
            35   40   45

Gly Ser Ser Gly Ser Ser
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Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys
1   5   10   15

Val Lys Lys Asp Glu Leu Cys Met Ser Met Ala Cys Val Cys Cys Glu
          20   25   30

Pro Leu Glu Cys Lys Tyr His Gly Tyr Phe Trp Leu Cys Gly Ser Gly
          35   40   45

Ser Ser Gly Ser Ser
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<213> Gymnea sylvestre

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1   5   10   15

Val Lys Lys Asp Glu Leu Cys Ala Val Trp Phe Asp Val Cys Cys Glu
          20   25   30

Pro Leu Glu Cys Thr Tyr Gln Ser Gly Tyr Tyr Trp Leu Cys Gly Ser

          35   40   45

Gly Ser Ser Gly Ser Ser

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<210> 33

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<400> 33

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1   5   10   15

Val Lys Lys Asp Glu Leu Cys Glu Pro Trp Tyr Trp Arg Cys Cys Glu

          20   25   30

Pro Leu Glu Cys Val Tyr Thr Ser Gly Tyr Tyr Tyr Ser Cys Gly Ser

          35   40   45

Gly Ser Ser Gly Ser Ser

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<400> 34

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1   5   10   15

Val Lys Lys Asp Glu Leu Cys Ala Arg Trp Asp Leu Val Cys Cys Glu

            20   25   30

Pro Leu Glu Cys Ile Tyr Thr Ser Glu Leu Tyr Ala Thr Cys Gly Ser

        35   40   45

Gly Ser Ser Gly Ser Ser

        50

<210> 35

<211> 54

<212> PRT

<213> Gymnea sylvestre

<400> 35

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1   5   10   15

Val Lys Lys Asp Glu Leu Cys Val Phe Tyr Phe Pro Asn Cys Cys Glu

           20   25   30

Pro Leu Glu Cys Arg Trp Val Asn Asp Asn Tyr Gly Trp Cys Gly Ser

      35   40   45

Gly Ser Ser Gly Ser Ser

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<210> 36

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<213> Gymnea sylvestre

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Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

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1              5                 10                15
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Val Lys Lys Asp Glu Leu Cys Met Ser Met Ala Cys Val Cys Cys Glu

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          20                25                30
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        35                40                45
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Ser Ser Gly Ser Ser

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<400> 37

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

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          20                25                30
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        35   40   45

Gly Ser Ser Gly Ser Ser

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<212> PRT

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<400> 38

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

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            20   25   30

Pro Leu Glu Cys Lys Trp Phe Asn Glu Asn Tyr Gly Ile Cys Gly Ser

        35   40   45

Gly Ser Ser Gly Ser Ser

    50

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Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1   5   10   15

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            20   25   30

Pro Leu Glu Cys Ile Tyr Thr Ser Glu Leu Tyr Ala Thr Cys Gly Ser

        35   40   45

Gly Ser Ser Gly Ser Ser

    50

<210> 40

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Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1   5   10   15

Val Lys Lys Asp Glu Leu Cys Ala Arg Trp Asp Leu Val Cys Cys Glu

            20   25   30

Pro Leu Glu Cys Leu Gly His Gly Leu Gly Tyr Ala Tyr Cys Gly Ser

        35   40   45

Gly Ser Ser Gly Ser Ser

    50

<210> 41

<211> 53

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<213> Gymnea sylvestre

<400> 41

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Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys
1               5                   10                  15


Val Lys Lys Asp Glu Leu Cys Met Trp Ser Arg Glu Val Cys Cys Glu
            20                  25                  30


Pro Leu Glu Cys Tyr Tyr Thr Gly Trp Tyr Trp Ala Cys Gly Ser Gly
        35                  40                  45


Ser Ser Gly Ser Ser
    50


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<212>  PRT

<213>  Gymnea sylvestre


<400>  42


Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys
1               5                   10                  15


Val Lys Lys Asp Glu Leu Cys Glu Leu Ala Val Asp Glu Cys Cys Glu
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        35   40   45

Gly Ser Ser Gly Ser Ser

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